



SEQUENCE LISTING

<110> Larenas, Edmund A.  
Goedegebuur, Frits  
Gualfetti, Peter  
Mitchinson, Colin

<120> Variant Humicola grisea CBH1.1

<130> GC794-2

<140> US 10/810,277  
<141> 2004-03-26

<150> US 60/459,734  
<151> 2003-04-01

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ggcgccggcc gctatggta ctgctgtct gagatggata tctggaaagc caacaacatg 780  
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gacttcaact cgtaccgcca gggcaacaag accttctacg gcaaggccat gaccgtcgac 960  
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<213> Humicola grisea

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cgtacctacc	tgtatggacgg	cgaggacaag	tatcagaccc	tcgagctct	cgccaacgag	180
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gccaacattg	agggctggac	cggtccacc	aacgacccca	acgcggcgc	gggcccgtat	420
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cctcaccctt	gcaccatcat	tggccagagc	cgctgcgagg	gcaactcg	cggtggcacc	540
tacagcaacg	agcgctacgc	cggtctgc	gaccccgatg	gctgcgactt	caactcg	600
cgccaggggca	acaagacctt	ctacggcaag	gcatgaccg	tcgacaccac	caagaagatc	660
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<211> 525

<212> PRT

<213> Humicola grisea

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Ser	Trp	Lys	Lys	Cys	Thr	Ala	Gly	Gly	Gln	Cys	Gln	Thr	Val	Gln	Ala	
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Ser	Ile	Thr	Leu	Asp	Ser	Asn	Trp	Arg	Trp	Trp	Thr	His	Gln	Val	Ser	Gly
							50		55				60			
Ser	Thr	Asn	Cys	Tyr	Thr	Gly	Asn	Lys	Trp	Asp	Thr	Ser	Ile	Cys	Thr	
							65		70				75			80
Asp	Ala	Lys	Ser	Cys	Ala	Gln	Asn	Cys	Cys	Val	Asp	Gly	Ala	Asp	Tyr	
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Thr	Ser	Thr	Tyr	Ile	Thr	Asn	Gly	Asp	Ser	Leu	Ser	Leu	Lys			
							100		105				110			
Phe	Val	Thr	Lys	Gly	Gln	His	Ser	Thr	Asn	Val	Gly	Ser	Arg	Thr	Tyr	
							115		120				125			
Leu	Met	Asp	Gly	Glu	Asp	Lys	Tyr	Gln	Thr	Phe	Glu	Leu	Leu	Gly	Asn	
							130		135				140			
Glu	Phe	Thr	Phe	Asp	Val	Asp	Val	Ser	Asn	Ile	Gly	Cys	Gly	Leu	Asn	
							145		150				155			160
Gly	Ala	Leu	Tyr	Phe	Val	Ser	Met	Asp	Ala	Asp	Gly	Gly	Leu	Ser	Arg	
							165		170				175			
Tyr	Pro	Gly	Asn	Lys	Ala	Gly	Ala	Lys	Tyr	Gly	Thr	Gly	Tyr	Cys	Asp	
							180		185				190			
Ala	Gln	Cys	Pro	Arg	Asp	Ile	Lys	Phe	Ile	Asn	Gly	Glu	Ala	Asn	Ile	
							195		200				205			
Glu	Gly	Trp	Thr	Gly	Ser	Thr	Asn	Asp	Pro	Asn	Ala	Gly	Ala	Gly	Arg	
							210		215				220			
Tyr	Gly	Thr	Cys	Cys	Ser	Glu	Met	Asp	Ile	Trp	Glu	Ala	Asn	Asn	Met	
							225		230				235			240

Ala Thr Ala Phe Thr Pro His Pro Cys Thr Ile Ile Gly Gln Ser Arg  
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 Cys Glu Gly Asp Ser Cys Gly Gly Thr Tyr Ser Asn Glu Arg Tyr Ala  
                  260                 265                 270  
 Gly Val Cys Asp Pro Asp Gly Cys Asp Phe Asn Ser Tyr Arg Gln Gly  
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 Asn Lys Thr Phe Tyr Gly Lys Gly Met Thr Val Asp Thr Thr Lys Lys  
                  290                 295                 300  
 Ile Thr Val Val Thr Gln Phe Leu Lys Asp Ala Asn Gly Asp Leu Gly  
                  305                 310                 315                 320  
 Glu Ile Lys Arg Phe Tyr Val Gln Asp Gly Lys Ile Ile Pro Asn Ser  
                  325                 330                 335  
 Glu Ser Thr Ile Pro Gly Val Glu Gly Asn Ser Ile Thr Gln Asp Trp  
                  340                 345                 350  
 Cys Asp Arg Gln Lys Val Ala Phe Gly Asp Ile Asp Asp Phe Asn Arg  
                  355                 360                 365  
 Lys Gly Gly Met Lys Gln Met Gly Lys Ala Leu Ala Gly Pro Met Val  
                  370                 375                 380  
 Leu Val Met Ser Ile Trp Asp Asp His Ala Ser Asn Met Leu Trp Leu  
                  385                 390                 395                 400  
 Asp Ser Thr Phe Pro Val Asp Ala Ala Gly Lys Pro Gly Ala Glu Arg  
                  405                 410                 415  
 Gly Ala Cys Pro Thr Thr Ser Gly Val Pro Ala Glu Val Glu Ala Glu  
                  420                 425                 430  
 Ala Pro Asn Ser Asn Val Val Phe Ser Asn Ile Arg Phe Gly Pro Ile  
                  435                 440                 445  
 Gly Ser Thr Val Ala Gly Leu Pro Gly Ala Gly Asn Gly Gly Asn Asn  
                  450                 455                 460  
 Gly Gly Asn Pro Pro Pro Pro Thr Thr Thr Thr Ser Ser Ala Pro Ala  
                  465                 470                 475                 480  
 Thr Thr Thr Ala Ser Ala Gly Pro Lys Ala Gly Arg Trp Gln Gln  
                  485                 490                 495  
 Cys Gly Gly Ile Gly Phe Thr Gly Pro Thr Gln Cys Glu Glu Pro Tyr  
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 <211> 507  
 <212> PRT  
 <213> Humicola grisea

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          20                 25                 30  
 Thr Leu Asp Ser Asn Trp Arg Trp Thr His Gln Val Ser Gly Ser Thr  
          35                 40                 45  
 Asn Cys Tyr Thr Gly Asn Lys Trp Asp Thr Ser Ile Cys Thr Asp Ala  
          50                 55                 60  
 Lys Ser Cys Ala Gln Asn Cys Cys Val Asp Gly Ala Asp Tyr Thr Ser  
          65                 70                 75                 80  
 Thr Tyr Gly Ile Thr Thr Asn Gly Asp Ser Leu Ser Leu Lys Phe Val  
          85                 90                 95  
 Thr Lys Gly Gln His Ser Thr Asn Val Gly Ser Arg Thr Tyr Leu Met  
          100                 105                 110  
 Asp Gly Glu Asp Lys Tyr Gln Thr Phe Glu Leu Leu Gly Asn Glu Phe  
          115                 120                 125  
 Thr Phe Asp Val Asp Val Ser Asn Ile Gly Cys Gly Leu Asn Gly Ala  
          130                 135                 140  
 Leu Tyr Phe Val Ser Met Asp Ala Asp Gly Gly Leu Ser Arg Tyr Pro

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Cys Pro Arg Asp Ile Lys Phe Ile Asn Gly Glu Ala Asn Ile Glu Gly			
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Trp Thr Gly Ser Thr Asn Asp Pro Asn Ala Gly Ala Gly Arg Tyr Gly			
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Thr Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Asn Met Ala Thr			
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Ala Phe Thr Pro His Pro Cys Thr Ile Ile Gly Gln Ser Arg Cys Glu			
225	230	235	240
Gly Asp Ser Cys Gly Gly Thr Tyr Ser Asn Glu Arg Tyr Ala Gly Val			
245	250	255	
Cys Asp Pro Asp Gly Cys Asp Phe Asn Ser Tyr Arg Gln Gly Asn Lys			
260	265	270	
Thr Phe Tyr Gly Lys Gly Met Thr Val Asp Thr Thr Lys Lys Ile Thr			
275	280	285	
Val Val Thr Gln Phe Leu Lys Asp Ala Asn Gly Asp Leu Gly Glu Ile			
290	295	300	
Lys Arg Phe Tyr Val Gln Asp Gly Lys Ile Ile Pro Asn Ser Glu Ser			
305	310	315	320
Thr Ile Pro Gly Val Glu Gly Asn Ser Ile Thr Gln Asp Trp Cys Asp			
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Arg Gln Lys Val Ala Phe Gly Asp Ile Asp Asp Phe Asn Arg Lys Gly			
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Gly Met Lys Gln Met Gly Lys Ala Leu Ala Gly Pro Met Val Leu Val			
355	360	365	
Met Ser Ile Trp Asp Asp His Ala Ser Asn Met Leu Trp Leu Asp Ser			
370	375	380	
Thr Phe Pro Val Asp Ala Ala Gly Lys Pro Gly Ala Glu Arg Gly Ala			
385	390	395	400
Cys Pro Thr Thr Ser Gly Val Pro Ala Glu Val Glu Ala Glu Ala Pro			
405	410	415	
Asn Ser Asn Val Val Phe Ser Asn Ile Arg Phe Gly Pro Ile Gly Ser			
420	425	430	
Thr Val Ala Gly Leu Pro Gly Ala Gly Asn Gly Gly Asn Asn Gly Gly			
435	440	445	
Asn Pro Pro Pro Thr Thr Ser Ser Ala Pro Ala Thr Thr			
450	455	460	
Thr Thr Ala Ser Ala Gly Pro Lys Ala Gly Arg Trp Gln Gln Cys Gly			
465	470	475	480
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Thr Lys Leu Asn Asp Trp Tyr Ser Gln Cys Leu			
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<211> 507  
<212> PRT  
<213> Humicola grisea

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Thr Leu Asp Ser Asn Trp Arg Trp Thr His Gln Val Ser Gly Ser Thr  
35 40 45  
Asn Cys Tyr Thr Gly Asn Lys Trp Asp Thr Ser Ile Cys Thr Asp Ala  
50 55 60  
Lys Ser Cys Ala Gln Asn Cys Cys Val Asp Gly Ala Asp Tyr Thr Ser  
65 70 75 80

Thr Tyr Gly Ile Thr Thr Asn Gly Asp Ser Leu Ser Leu Lys Phe Val  
 85 90 95  
 Thr Lys Gly Gln His Ser Thr Asn Val Gly Ser Arg Thr Tyr Leu Met  
 100 105 110  
 Asp Gly Glu Asp Lys Tyr Gln Thr Phe Glu Leu Leu Gly Asn Glu Phe  
 115 120 125  
 Thr Phe Asp Val Asp Val Ser Asn Ile Gly Cys Gly Leu Asn Gly Ala  
 130 135 140  
 Leu Tyr Phe Val Ser Met Asp Ala Asp Gly Gly Leu Ser Arg Tyr Pro  
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 Gly Asn Lys Ala Gly Ala Lys Tyr Gly Thr Gly Tyr Cys Asp Ala Gln  
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 Cys Pro Arg Asp Ile Lys Phe Ile Asn Gly Glu Ala Asn Ile Glu Gly  
 180 185 190  
 Trp Thr Gly Ser Thr Asn Asp Pro Asn Ala Gly Ala Gly Arg Tyr Gly  
 195 200 205  
 Thr Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Asn Met Ala Thr  
 210 215 220  
 Ala Phe Thr Pro His Pro Cys Thr Ile Ile Gly Gln Ser Arg Cys Glu  
 225 230 235 240  
 Gly Asp Ser Cys Gly Gly Thr Tyr Ser Asn Glu Arg Tyr Ala Gly Val  
 245 250 255  
 Cys Asp Pro Asp Gly Cys Asp Phe Asn Ser Tyr Arg Gln Gly Asn Lys  
 260 265 270  
 Thr Phe Tyr Gly Lys Gly Met Thr Val Asp Thr Thr Lys Lys Ile Thr  
 275 280 285  
 Val Val Thr Gln Phe Leu Lys Asp Ala Asn Gly Asp Leu Gly Glu Ile  
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 Lys Arg Phe Tyr Val Gln Asp Gly Lys Ile Ile Pro Asn Ser Glu Ser  
 305 310 315 320  
 Thr Ile Pro Gly Val Glu Gly Asn Ser Ile Thr Gln Asp Trp Cys Asp  
 325 330 335  
 Arg Gln Lys Val Ala Phe Gly Asp Ile Asp Asp Phe Asn Arg Lys Gly  
 340 345 350  
 Gly Met Lys Gln Met Gly Lys Ala Leu Ala Gly Pro Met Val Leu Val  
 355 360 365  
 Met Ser Ile Trp Asp Asp His Ala Ser Asn Met Leu Trp Leu Asp Ser  
 370 375 380  
 Thr Phe Pro Val Asp Ala Ala Gly Lys Pro Gly Ala Glu Arg Gly Ala  
 385 390 395 400  
 Cys Pro Thr Thr Ser Gly Val Pro Ala Glu Val Glu Ala Glu Ala Pro  
 405 410 415  
 Asn Ser Asn Val Val Phe Ser Asn Ile Arg Phe Gly Pro Ile Gly Ser  
 420 425 430  
 Thr Val Ala Gly Leu Pro Gly Ala Gly Asn Gly Gly Asn Asn Gly Gly  
 435 440 445  
 Asn Pro Pro Pro Thr Thr Ser Ser Ala Pro Ala Thr Thr  
 450 455 460  
 Thr Thr Ala Ser Ala Gly Pro Lys Ala Gly Arg Trp Gln Gln Cys Gly  
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 Thr Lys Leu Asn Asp Trp Tyr Ser Gln Cys Leu  
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<210> 6  
 <211> 507  
 <212> PRT  
 <213> Humicola grisea

<400> 6  
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Thr	Leu	Asp	Ser	Asn	Trp	Arg	Trp	Thr	His	Gln	Val	Ser	Gly	Ser	Thr
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Asn	Cys	Tyr	Thr	Gly	Asn	Lys	Trp	Asp	Thr	Ser	Ile	Cys	Thr	Asp	Ala
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Lys	Ser	Cys	Ala	Gln	Asn	Cys	Cys	Val	Asp	Gly	Ala	Asp	Tyr	Thr	Ser
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Thr	Tyr	Gly	Ile	Thr	Thr	Asn	Gly	Asp	Ser	Leu	Ser	Leu	Lys	Phe	Val
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Thr	Lys	Gly	Gln	Tyr	Ser	Thr	Asn	Val	Gly	Ser	Arg	Thr	Tyr	Leu	Met
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Asp	Gly	Glu	Asp	Lys	Tyr	Gln	Thr	Phe	Glu	Leu	Leu	Gly	Asn	Glu	Phe
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Thr	Phe	Asp	Val	Asp	Val	Ser	Asn	Ile	Gly	Cys	Gly	Leu	Asn	Gly	Ala
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Leu	Tyr	Phe	Val	Ser	Met	Asp	Ala	Asp	Gly	Gly	Leu	Ser	Arg	Tyr	Pro
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Gly	Asn	Lys	Ala	Gly	Ala	Lys	Tyr	Gly	Thr	Gly	Tyr	Cys	Asp	Ala	Gln
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Cys	Pro	Arg	Asp	Ile	Lys	Phe	Ile	Asn	Gly	Glu	Ala	Asn	Ile	Glu	Gly
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Trp	Thr	Gly	Ser	Thr	Asn	Asp	Pro	Asn	Ala	Gly	Ala	Gly	Arg	Tyr	Gly
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Thr	Cys	Cys	Ser	Glu	Met	Asp	Ile	Trp	Glu	Ala	Asn	Asn	Met	Ala	Thr
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Ala	Phe	Thr	Pro	His	Pro	Cys	Thr	Ile	Ile	Gly	Gln	Ser	Arg	Cys	Glu
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Gly	Asp	Ser	Cys	Gly	Gly	Thr	Tyr	Ser	Asn	Glu	Arg	Tyr	Ala	Gly	Val
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Cys	Asp	Pro	Asp	Gly	Cys	Asp	Phe	Asn	Ser	Tyr	Arg	Gln	Gly	Asn	Lys
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Thr	Phe	Tyr	Gly	Lys	Gly	Met	Thr	Val	Asp	Thr	Thr	Lys	Lys	Ile	Thr
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Val	Val	Thr	Gln	Phe	Leu	Lys	Asp	Ala	Asn	Gly	Asp	Leu	Gly	Glu	Ile
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Lys	Arg	Phe	Tyr	Val	Gln	Asp	Gly	Lys	Ile	Ile	Pro	Asn	Ser	Glu	Ser
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Thr	Ile	Pro	Gly	Val	Glu	Gly	Asn	Ser	Ile	Thr	Gln	Asp	Trp	Cys	Asp
			325				330				335				
Arg	Gln	Lys	Val	Ala	Phe	Gly	Asp	Ile	Asp	Asp	Phe	Asn	Arg	Lys	Gly
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Gly	Met	Lys	Gln	Met	Gly	Lys	Ala	Leu	Ala	Gly	Pro	Met	Val	Leu	Val
			355				360				365				
Met	Ser	Ile	Trp	Asp	Asp	His	Ala	Ser	Asn	Met	Leu	Trp	Leu	Asp	Ser
			370				375				380				
Thr	Phe	Pro	Val	Asp	Ala	Ala	Gly	Lys	Pro	Gly	Ala	Glu	Arg	Gly	Ala
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Cys	Pro	Thr	Thr	Ser	Gly	Val	Pro	Ala	Glu	Val	Glu	Ala	Glu	Ala	Pro
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Asn	Ser	Asn	Val	Val	Phe	Ser	Asn	Ile	Arg	Phe	Gly	Pro	Ile	Gly	Ser
			420				425				430				
Thr	Val	Ala	Gly	Leu	Pro	Gly	Ala	Gly	Asn	Gly	Gly	Asn	Asn	Gly	Gly
			435				440				445				
Asn	Pro	Pro	Pro	Pro	Thr	Thr	Thr	Ser	Ser	Ala	Pro	Ala	Thr	Thr	
			450				455				460				
Thr	Thr	Ala	Ser	Ala	Gly	Pro	Lys	Ala	Gly	Arg	Trp	Gln	Gln	Cys	Gly
			465				470				475			480	
Gly	Ile	Gly	Phe	Thr	Gly	Pro	Thr	Gln	Cys	Glu	Glu	Pro	Tyr	Thr	Cys
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Thr	Lys	Leu	Asn	Asp	Trp	Tyr	Ser	Gln	Cys	Leu					

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505

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 <211> 1662  
 <212> DNA  
 <213> *Scytalidium thermophilium*

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caggtgtctg	gctccaccaa	ctgctacacg	ggcaacgagt	gggattctag	catctgcact	240
gatgccaagt	cgtgcgtca	gaactgctgc	gtcgatgggt	ctgactacac	cagcacctat	300
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ctcctcgcc	acgagttcac	cttcgatgtc	gatgtctcca	acatcggtcg	cggtctcaac	540
ggccccctgt	acttcgtctc	catggacgccc	gatgggtggc	tcagggctca	tcctggcaac	600
aaggctggtg	ccaagtacgg	taccggctac	tgcgatgctc	agtggccccc	tgacatcaag	660
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tcgtgcgttg	gcacccatag	caacgaccgc	tacggccggc	tctgggatcc	cgatggctgc	900
gacttcaacg	cgtatcgcca	gggcaacaa	accttctacg	gcaagggcat	gaccgtcgac	960
accaccaaga	agtcaccgt	cgtaaccccg	ttcctcaagg	acgccaacgg	cgatctcgcc	1020
gagatcaacg	gcttctacgt	ccaggatggg	aaagatcatcc	ccaaactccga	gtccaccatc	1080
ccccgcgtcg	aggcaactc	catcaccatc	gattgggtcg	accggccagaa	gtttgccttt	1140
ggcgacattg	acgacttcaa	ccgcaaggcc	ggcatgaacg	agatgggcaa	ggccctcgcc	1200
ggcccccattgg	tcctggatcat	gtccatctgg	gatgaccacg	cctccaacat	gctctggctc	1260
gactcgacct	tcctgtcga	tgcgcgtggc	aagccggcg	ccgagcgcgg	tgcctggccg	1320
accacctcgg	gtgtccctgc	tgaggttgag	gccgaggccc	ccaaacagcaa	cgtcgtcttc	1380
tccaacatcc	gctcggtccc	catcggtctc	accgttgccg	gccttcccg	cgatggcgcc	1440
aacaacggcg	gcaacaccac	cgtaacggcc	ccggccagca	ccaccaccac	ctctggccagc	1500
agcagcacca	cctcggtcc	tgccaccacc	accaccgcca	gctggggccc	caaggctggc	1560
cgctggcagc	agtgcggcgg	catcggttcc	actggcccg	cccagtgcga	ggagccctac	1620
acttgcacca	agtcacacga	ctggtaactct	cagtgcctgt	aa		1662

<210> 8  
 <211> 1602  
 <212> DNA  
 <213> *Scytalidium thermophilium*

<400> 8

atgcgtaccc	ccaagttcgc	caccctcgcc	gcccttgtgg	cctcgccgc	cgcccagcag	60
gcgtgcagcc	tcaccaccga	gaggcacccct	tccctctct	ggaagaagtgc	caccgcggc	120
ggccagtgcc	agaccgtcca	ggcttcatc	actctcgact	ccaactggcg	ctggactcac	180
caggtgtctg	gctccaccaa	ctgctacacg	ggcaacgagt	gggattctag	catctgcact	240
gatgccaagt	cgtgcgtca	gaactgctgc	gtcgatgggt	ctgactacac	cagcacctat	300
ggcatcacca	ccaacggtga	ttccctgagc	ctcaagttcg	tcaccaaggg	ccagtaactcg	360
accaacgtcg	gctcggtac	ctacctgatg	gacggcgagg	acaagtatca	gaccctcgag	420
ctcctcgcc	acgagttcac	cttcgatgtc	gatgtctcca	acatcggtcg	cggtctcaac	480
ggcgccctgt	acttcgtctc	catggacgccc	gatgggtggc	tcagggctca	tcctggcaac	540
aaggctgttg	ccaagtacgg	taccggctac	tgcgatgctc	agtggccccc	tgacatcaag	600
ttcatcaacg	gcgaggccaa	cattgagggc	tggaccggct	ccaccaacga	ccccaacgccc	660
ggcgccggcc	gctatggtac	ctgctgtct	gagatggata	tctgggaggg	caacaacatg	720
gtactgcct	tcactcctca	cccttgact	atcattggcc	agagccgctg	cgagggcgac	780
tcgtgcgttg	gcacccatag	caacgaccgc	tacggccggc	tctgcaccc	cgatggctgc	840
gacttcaacg	cgtatcgcca	gggcaacaa	accttctacg	gcaagggcat	gaccgtcgac	900
accaccaaga	agtcaccgt	cgtaaccccg	ttcctcaagg	acgccaacgg	cgatctcgcc	960
gagatcaacg	gcttctacgt	ccaggatggg	aaagatcatcc	ccaaactccga	gtccaccatc	1020
ccccgcgtcg	aggcaactc	catcaccatc	gattgggtcg	accggccagaa	gtttgccttt	1080
ggcgacattg	acgacttcaa	ccgcaaggcc	ggcatgaacg	agatgggcaa	ggccctcgcc	1140

ggccccatgg	tcctggtcat	gtccatctgg	gatgaccacg	cctccaacat	gctctggctc	1200
'gactcgacct	tccctgtcga	tgccgctggc	aagcccggcg	ccgagcgcgg	tgcctgccc	1260
accacacccgg	gtgtccctgc	tgaggttgag	gccgaggccc	ccaacagcaa	cgtcgcttc	1320
tccaaacatcc	gcttcggccc	catcgctcg	accgttgccg	gccttcccag	cgatggcggc	1380
aacaacggcg	gcaacaccac	cgtccagccc	ccgcccagca	ccaccaccac	ctctgccagc	1440
agcagcacca	cctcggtc	tgccaccacc	accaccgcca	ggcgtggccc	caaggctggc	1500
cgctggcagc	agtgcggcgg	catcgcttc	actggccca	cccagtgcga	ggagccctac	1560
acttgcacca	agctcaacga	ctggta	cagtgcctgt	aa		1602

<210> 9

<211> 533

<212> PRT

<213> *Scytalidium thermophilium*

<400> 9

Met	Arg	Thr	Ala	Lys	Phe	Ala	Thr	Leu	Ala	Ala	Leu	Val	Ala	Ser	Ala
1															
														15	
Ala	Ala	Gln	Gln	Ala	Cys	Ser	Leu	Thr	Thr	Glu	Arg	His	Pro	Ser	Leu
														30	
Ser	Trp	Lys	Lys	Cys	Thr	Ala	Gly	Gly	Gln	Cys	Gln	Thr	Val	Gln	Ala
														45	
Ser	Ile	Thr	Leu	Asp	Ser	Asn	Trp	Arg	Trp	Thr	His	Gln	Val	Ser	Gly
														50	
														55	
														60	
Ser	Thr	Asn	Cys	Tyr	Thr	Gly	Asn	Glu	Trp	Asp	Ser	Ser	Ile	Cys	Thr
														65	
														70	
														75	
														80	
Asp	Ala	Lys	Ser	Cys	Ala	Gln	Asn	Cys	Cys	Val	Asp	Gly	Ala	Asp	Tyr
														85	
														90	
														95	
Thr	Ser	Thr	Tyr	Ile	Thr	Thr	Asn	Gly	Asp	Ser	Leu	Ser	Leu	Lys	
														100	
														105	
														110	
Phe	Val	Thr	Lys	Gly	Gln	Tyr	Ser	Thr	Asn	Val	Gly	Ser	Arg	Thr	Tyr
														115	
														120	
														125	
Leu	Met	Asp	Gly	Glu	Asp	Lys	Tyr	Gln	Thr	Phe	Glu	Leu	Leu	Gly	Asn
														130	
														135	
														140	
Glu	Phe	Thr	Phe	Asp	Val	Asp	Val	Ser	Asn	Ile	Gly	Cys	Gly	Leu	Asn
														145	
														150	
														155	
Gly	Ala	Leu	Tyr	Phe	Val	Ser	Met	Asp	Ala	Asp	Gly	Gly	Leu	Ser	Arg
														165	
														170	
														175	
Tyr	Pro	Gly	Asn	Lys	Ala	Gly	Ala	Lys	Tyr	Gly	Thr	Gly	Tyr	Cys	Asp
														180	
														185	
														190	
Ala	Gln	Cys	Pro	Arg	Asp	Ile	Lys	Phe	Ile	Asn	Gly	Glu	Ala	Asn	Ile
														195	
														200	
														205	
Glu	Gly	Trp	Thr	Gly	Ser	Thr	Asn	Asp	Pro	Asn	Ala	Gly	Ala	Gly	Arg
														210	
														215	
														220	
Tyr	Gly	Thr	Cys	Cys	Ser	Glu	Met	Asp	Ile	Trp	Glu	Ala	Asn	Asn	Met
														225	
														230	
														235	
Ala	Thr	Ala	Phe	Thr	Pro	His	Pro	Cys	Thr	Ile	Ile	Gly	Gln	Ser	Arg
														245	
														250	
														255	
Cys	Glu	Gly	Asp	Ser	Cys	Gly	Gly	Thr	Tyr	Ser	Asn	Asp	Arg	Tyr	Ala
														260	
														265	
														270	
Gly	Val	Cys	Asp	Pro	Asp	Gly	Cys	Asp	Phe	Asn	Ala	Tyr	Arg	Gln	Gly
														275	
														280	
														285	
Asn	Lys	Thr	Phe	Tyr	Gly	Lys	Gly	Met	Thr	Val	Asp	Thr	Thr	Lys	Lys
														290	
														295	
														300	
Leu	Thr	Val	Val	Thr	Gln	Phe	Leu	Lys	Asp	Ala	Asn	Gly	Asp	Leu	Gly
														305	
														310	
														315	
Glu	Ile	Lys	Arg	Phe	Tyr	Val	Gln	Asp	Gly	Lys	Ile	Ile	Pro	Asn	Ser
														325	
														330	
														335	
Glu	Ser	Thr	Ile	Pro	Gly	Val	Glu	Gly	Asn	Ser	Ile	Thr	Gln	Asp	Trp
														340	
														345	
														350	
Cys	Asp	Arg	Gln	Lys	Val	Ala	Phe	Gly	Asp	Ile	Asp	Asp	Phe	Asn	Arg
														355	
														360	
														365	
Lys	Gly	Gly	Met	Lys	Gln	Met	Gly	Lys	Ala	Leu	Ala	Gly	Pro	Met	Val

370	375	380	
Leu Val Met Ser Ile Trp Asp Asp His Ala Ser Asn Met	Leu Trp Leu		
385	390	395	400
Asp Ser Thr Phe Pro Val Asp Ala Ala Gly Lys Pro Gly Ala Glu Arg			
405	410	415	
Gly Ala Cys Pro Thr Thr Ser Gly Val Pro Ala Glu Val Glu Ala Glu			
420	425	430	
Ala Pro Asn Ser Asn Val Val Phe Ser Asn Ile Arg Phe Gly Pro Ile			
435	440	445	
Gly Ser Thr Val Ala Gly Leu Pro Ser Asp Gly Gly Asn Asn Gly Gly			
450	455	460	
Asn Thr Thr Val Gln Pro Pro Ser Thr Thr Thr Ser Ala Ser			
465	470	475	480
Ser Ser Thr Thr Ser Ala Pro Ala Thr Thr Thr Ala Ser Ala Gly			
485	490	495	
Pro Lys Ala Gly Arg Trp Gln Gln Cys Gly Gly Ile Gly Phe Thr Gly			
500	505	510	
Pro Thr Gln Cys Glu Glu Pro Tyr Thr Cys Thr Lys Leu Asn Asp Trp			
515	520	525	
Tyr Ser Gln Cys Leu			
530			

<210> 10

<211> 497

<212> PRT

<213> Hypocrea jecorina

<400> 10

Gln Ser Ala Cys Thr Leu Gln Ser Glu Thr His Pro Pro Leu Thr Trp			
1	5	10	15
Gln Lys Cys Ser Ser Gly Gly Thr Cys Thr Gln Gln Thr Gly Ser Val			
20	25	30	
Val Ile Asp Ala Asn Trp Arg Trp Thr His Ala Thr Asn Ser Ser Thr			
35	40	45	
Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Thr Leu Cys Pro Asp Asn			
50	55	60	
Glu Thr Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser			
65	70	75	80
Thr Tyr Gly Val Thr Thr Ser Gly Asn Ser Leu Ser Ile Gly Phe Val			
85	90	95	
Thr Gln Ser Ala Gln Lys Asn Val Gly Ala Arg Leu Tyr Leu Met Ala			
100	105	110	
Ser Asp Thr Thr Tyr Gln Glu Phe Thr Leu Leu Gly Asn Glu Phe Ser			
115	120	125	
Phe Asp Val Asp Val Ser Gln Leu Pro Cys Gly Leu Asn Gly Ala Leu			
130	135	140	
Tyr Phe Val Ser Met Asp Ala Asp Gly Gly Val Ser Lys Tyr Pro Thr			
145	150	155	160
Asn Thr Ala Gly Ala Lys Tyr Gly Thr Gly Tyr Cys Asp Ser Gln Cys			
165	170	175	
Pro Arg Asp Leu Lys Phe Ile Asn Gly Gln Ala Asn Val Glu Gly Trp			
180	185	190	
Glu Pro Ser Ser Asn Asn Ala Asn Thr Gly Ile Gly Gly His Gly Ser			
195	200	205	
Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Ser Ile Ser Glu Ala			
210	215	220	
Leu Thr Pro His Pro Cys Thr Thr Val Gly Gln Glu Ile Cys Glu Gly			
225	230	235	240
Asp Gly Cys Gly Gly Thr Tyr Ser Asp Asn Arg Tyr Gly Gly Thr Cys			
245	250	255	
Asp Pro Asp Gly Cys Asp Trp Asn Pro Tyr Arg Leu Gly Asn Thr Ser			
260	265	270	

Phe Tyr Gly Pro Gly Ser Ser Phe Thr Leu Asp Thr Thr Lys Lys Leu  
 275 280 285  
 Thr Val Val Thr Gln Phe Glu Thr Ser Gly Ala Ile Asn Arg Tyr Tyr  
 290 295 300  
 Val Gln Asn Gly Val Thr Phe Gln Gln Pro Asn Ala Glu Leu Gly Ser  
 305 310 315 320  
 Tyr Ser Gly Asn Glu Leu Asn Asp Asp Tyr Cys Thr Ala Glu Glu Ala  
 325 330 335  
 Glu Phe Gly Gly Ser Ser Phe Ser Asp Lys Gly Gly Leu Thr Gln Phe  
 340 345 350  
 Lys Lys Ala Thr Ser Gly Gly Met Val Leu Val Met Ser Leu Trp Asp  
 355 360 365  
 Asp Tyr Tyr Ala Asn Met Leu Trp Leu Asp Ser Thr Tyr Pro Thr Asn  
 370 375 380  
 Glu Thr Ser Ser Thr Pro Gly Ala Val Arg Gly Ser Cys Ser Thr Ser  
 385 390 395 400  
 Ser Gly Val Pro Ala Gln Val Glu Ser Gln Ser Pro Asn Ala Lys Val  
 405 410 415  
 Thr Phe Ser Asn Ile Lys Phe Gly Pro Ile Gly Ser Thr Gly Asn Pro  
 420 425 430  
 Ser Gly Gly Asn Pro Pro Gly Gly Asn Pro Pro Gly Thr Thr Thr Thr  
 435 440 445  
 Arg Arg Pro Ala Thr Thr Thr Gly Ser Ser Pro Gly Pro Thr Gln Ser  
 450 455 460  
 His Tyr Gly Gln Cys Gly Gly Ile Gly Tyr Ser Gly Pro Thr Val Cys  
 465 470 475 480  
 Ala Ser Gly Thr Thr Cys Gln Val Leu Asn Pro Tyr Tyr Ser Gln Cys  
 485 490 495  
 Leu

<210> 11  
 <211> 515  
 <212> PRT  
 <213> *Scy whole* *thermophilum*

<400> 11  
 Gln Gln Ala Cys Ser Leu Thr Thr Glu Arg His Pro Ser Leu Ser Trp  
 1 5 10 15  
 Lys Lys Cys Thr Ala Gly Gly Gln Cys Gln Thr Val Gln Ala Ser Ile  
 20 25 30  
 Thr Leu Asp Ser Asn Trp Arg Trp Thr His Gln Val Ser Gly Ser Thr  
 35 40 45  
 Asn Cys Tyr Thr Gly Asn Glu Trp Asp Ser Ser Ile Cys Thr Asp Ala  
 50 55 60  
 Lys Ser Cys Ala Gln Asn Cys Cys Val Asp Gly Ala Asp Tyr Thr Ser  
 65 70 75 80  
 Thr Tyr Gly Ile Thr Thr Asn Gly Asp Ser Leu Ser Leu Lys Phe Val  
 85 90 95  
 Thr Lys Gly Gln Tyr Ser Thr Asn Val Gly Ser Arg Thr Tyr Leu Met  
 100 105 110  
 Asp Gly Glu Asp Lys Tyr Gln Thr Phe Glu Leu Leu Gly Asn Glu Phe  
 115 120 125  
 Thr Phe Asp Val Asp Val Ser Asn Ile Gly Cys Gly Leu Asn Gly Ala  
 130 135 140  
 Leu Tyr Phe Val Ser Met Asp Ala Asp Gly Gly Leu Ser Arg Tyr Pro  
 145 150 155 160  
 Gly Asn Lys Ala Gly Ala Lys Tyr Gly Thr Gly Tyr Cys Asp Ala Gln  
 165 170 175  
 Cys Pro Arg Asp Ile Lys Phe Ile Asn Gly Glu Ala Asn Ile Glu Gly  
 180 185 190  
 Trp Thr Gly Ser Thr Asn Asp Pro Asn Ala Gly Ala Gly Arg Tyr Gly

195	200	205
Thr Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Asn Met Ala Thr		
210	215	220
Ala Phe Thr Pro His Pro Cys Thr Ile Ile Gly Gln Ser Arg Cys Glu		
225	230	235
Gly Asp Ser Cys Gly Gly Thr Tyr Ser Asn Asp Arg Tyr Ala Gly Val		
245	250	255
Cys Asp Pro Asp Gly Cys Asp Phe Asn Ala Tyr Arg Gln Gly Asn Lys		
260	265	270
Thr Phe Tyr Gly Lys Gly Met Thr Val Asp Thr Thr Lys Lys Leu Thr		
275	280	285
Val Val Thr Gln Phe Leu Lys Asp Ala Asn Gly Asp Leu Gly Glu Ile		
290	295	300
Lys Arg Phe Tyr Val Gln Asp Gly Lys Ile Ile Pro Asn Ser Glu Ser		
305	310	315
Thr Ile Pro Gly Val Glu Gly Asn Ser Ile Thr Gln Asp Trp Cys Asp		
325	330	335
Arg Gln Lys Val Ala Phe Gly Asp Ile Asp Asp Phe Asn Arg Lys Gly		
340	345	350
Gly Met Lys Gln Met Gly Lys Ala Leu Ala Gly Pro Met Val Leu Val		
355	360	365
Met Ser Ile Trp Asp Asp His Ala Ser Asn Met Leu Trp Leu Asp Ser		
370	375	380
Thr Phe Pro Val Asp Ala Ala Gly Lys Pro Gly Ala Glu Arg Gly Ala		
385	390	395
Cys Pro Thr Thr Ser Gly Val Pro Ala Glu Val Glu Ala Glu Ala Pro		
405	410	415
Asn Ser Asn Val Val Phe Ser Asn Ile Arg Phe Gly Pro Ile Gly Ser		
420	425	430
Thr Val Ala Gly Leu Pro Ser Asp Gly Gly Asn Asn Gly Gly Asn Thr		
435	440	445
Thr Val Gln Pro Pro Pro Ser Thr Thr Thr Ser Ala Ser Ser Ser		
450	455	460
Thr Thr Ser Ala Pro Ala Thr Thr Thr Ala Ser Ala Gly Pro Lys		
465	470	475
Ala Gly Arg Trp Gln Gln Cys Gly Gly Ile Gly Phe Thr Gly Pro Thr		
485	490	495
Gln Cys Glu Glu Pro Tyr Thr Cys Thr Lys Leu Asn Asp Trp Tyr Ser		
500	505	510
Gln Cys Leu		
515		

<210> 12  
 <211> 507  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> consensus sequence

<400> 12  
 Gln Gln Ala Cys Ser Leu Thr Thr Glu Arg His Pro Ser Leu Ser Trp  
 1 5 10 15  
 Lys Lys Cys Thr Ala Gly Gly Gln Cys Gln Thr Val Gln Ala Ser Ile  
 20 25 30  
 Thr Leu Asp Ser Asn Trp Arg Trp Thr His Gln Val Ser Gly Ser Thr  
 35 40 45  
 Asn Cys Tyr Thr Gly Asn Lys Trp Asp Ser Ser Ile Cys Thr Asp Ala  
 50 55 60  
 Lys Ser Cys Ala Gln Asn Cys Cys Val Asp Gly Ala Asp Tyr Thr Ser  
 65 70 75 80  
 Thr Tyr Gly Ile Thr Thr Asn Gly Asp Ser Leu Ser Leu Lys Phe Val

85	90	95
Thr Lys Gly Gln His Ser Thr Asn Val	Gly Ser Arg Thr Tyr	Leu Met
100	105	110
Asp Gly Glu Asp Lys Tyr Gln Thr Phe	Glu Leu Leu Gly Asn Glu Phe	
115	120	125
Thr Phe Asp Val Asp Val Ser Asn Ile	Gly Cys Gly Leu Asn Gly Ala	
130	135	140
Leu Tyr Phe Val Ser Met Asp Ala Asp	Gly Gly Leu Ser Arg Tyr Pro	
145	150	160
Gly Asn Lys Ala Gly Ala Lys Tyr Gly	Thr Gly Tyr Cys Asp Ala Gln	
165	170	175
Cys Pro Arg Asp Ile Lys Phe Ile Asn	Gly Glu Ala Asn Ile Glu Gly	
180	185	190
Trp Thr Gly Ser Thr Asn Asp Pro Asn	Ala Gly Ala Gly Arg Tyr Gly	
195	200	205
Thr Cys Cys Ser Glu Met Asp Ile Trp	Glu Ala Asn Asn Met Ala Thr	
210	215	220
Ala Phe Thr Pro His Pro Cys Thr Ile	Ile Gly Gln Ser Arg Cys Glu	
225	230	240
Gly Asp Ser Cys Gly Gly Thr Tyr Ser	Asn Glu Arg Tyr Ala Gly Val	
245	250	255
Cys Asp Pro Asp Gly Cys Asp Phe Asn	Ser Tyr Arg Gln Gly Asn Lys	
260	265	270
Thr Phe Tyr Gly Lys Gly Met Thr Val	Asp Thr Thr Lys Lys Ile Thr	
275	280	285
Val Val Thr Gln Phe Leu Lys Asp Ala	Asn Gly Asp Leu Gly Glu Ile	
290	295	300
Lys Arg Phe Tyr Val Gln Asp Gly Lys	Ile Ile Pro Asn Ser Glu Ser	
305	310	320
Thr Ile Pro Gly Val Glu Gly Asn Ser	Ile Thr Gln Asp Trp Cys Asp	
325	330	335
Arg Gln Lys Val Ala Phe Gly Asp Ile	Asp Asp Phe Asn Arg Lys Gly	
340	345	350
Gly Met Lys Gln Met Gly Lys Ala Leu	Ala Gly Pro Met Val Leu Val	
355	360	365
Met Ser Ile Trp Asp Asp His Ala Ser	Asn Met Leu Trp Leu Asp Ser	
370	375	380
Thr Phe Pro Val Asp Ala Ala Gly Lys	Pro Gly Ala Glu Arg Gly Ala	
385	390	400
Cys Pro Thr Thr Ser Gly Val Pro Ala	Glu Val Glu Ala Glu Ala Pro	
405	410	415
Asn Ser Asn Val Val Phe Ser Asn Ile	Arg Phe Gly Pro Ile Gly Ser	
420	425	430
Thr Val Ala Gly Leu Pro Gly Ala Gly	Asn Gly Asn Asn Gly Gly	
435	440	445
Asn Pro Pro Pro Thr Thr Ser Ser Ala	Pro Ala Thr Thr	
450	455	460
Thr Thr Ala Ser Ala Gly Pro Lys Ala	Gly Arg Trp Gln Gln Cys Gly	
465	470	480
Gly Ile Gly Phe Thr Gly Pro Thr Gln	Cys Glu Glu Pro Tyr Thr Cys	
485	490	495
Thr Lys Leu Asn Asp Trp Tyr Ser Gln	Cys Leu	
500	505	

<210> 13

<211> 507

<212> PRT

<213> Artificial Sequence

<220>

<223> consensus sequence

<400> 13

Gln Gln Ala Cys Ser Leu Thr Thr Glu Arg His Pro Ser Leu Ser Trp  
1 5 10 15  
Lys Lys Cys Thr Ala Gly Gly Gln Cys Gln Thr Val Gln Ala Ser Ile  
20 25 30  
Thr Leu Asp Ser Asn Trp Arg Trp Thr His Gln Val Ser Gly Ser Thr  
35 40 45  
Asn Cys Tyr Thr Gly Asn Lys Trp Asp Thr Ser Ile Cys Thr Asp Ala  
50 55 60  
Lys Ser Cys Ala Gln Asn Cys Cys Val Asp Gly Ala Asp Tyr Thr Ser  
65 70 75 80  
Thr Tyr Gly Ile Thr Thr Asn Gly Asp Ser Leu Ser Leu Lys Phe Val  
85 90 95  
Thr Lys Gly Gln His Ser Thr Asn Val Gly Ser Arg Thr Tyr Leu Met  
100 105 110  
Asp Gly Glu Asp Lys Tyr Gln Thr Phe Glu Leu Leu Gly Asn Glu Phe  
115 120 125  
Thr Phe Asp Val Asp Val Ser Asn Ile Gly Cys Gly Leu Asn Gly Ala  
130 135 140  
Leu Tyr Phe Val Ser Met Asp Ala Asp Gly Gly Leu Ser Arg Tyr Pro  
145 150 155 160  
Gly Asn Lys Ala Gly Ala Lys Tyr Gly Thr Gly Tyr Cys Asp Ala Gln  
165 170 175  
Cys Pro Arg Asp Ile Lys Phe Ile Asn Gly Glu Ala Asn Ile Glu Gly  
180 185 190  
Trp Thr Gly Ser Thr Asn Asp Pro Asn Ala Gly Ala Gly Arg Tyr Gly  
195 200 205  
Thr Cys Cys Ser Glu Met Asp Ile Trp Glu Ala Asn Asn Met Ala Thr  
210 215 220  
Ala Phe Thr Pro His Pro Cys Thr Ile Ile Gly Gln Ser Arg Cys Glu  
225 230 235 240  
Gly Asp Ser Cys Gly Gly Thr Tyr Ser Asn Glu Arg Tyr Ala Gly Val  
245 250 255  
Cys Asp Pro Asp Gly Cys Asp Phe Asn Ser Tyr Arg Gln Gly Asn Lys  
260 265 270  
Thr Phe Tyr Gly Lys Gly Met Thr Val Asp Thr Thr Lys Lys Ile Thr  
275 280 285  
Val Val Thr Gln Phe Leu Lys Asp Ala Asn Gly Asp Leu Gly Glu Ile  
290 295 300  
Lys Arg Phe Tyr Val Gln Asp Gly Lys Ile Ile Pro Asn Ser Glu Ser  
305 310 315 320  
Thr Ile Pro Gly Val Glu Gly Asn Ser Ile Thr Gln Asp Trp Cys Asp  
325 330 335  
Arg Gln Lys Val Ala Phe Gly Asp Ile Asp Asp Phe Asn Arg Lys Gly  
340 345 350  
Gly Met Lys Gln Met Gly Lys Ala Leu Ala Gly Pro Met Val Leu Val  
355 360 365  
Met Ser Ile Trp Asp Asp His Ala Ser Asn Met Leu Trp Leu Asp Ser  
370 375 380  
Thr Phe Pro Val Asp Ala Ala Gly Lys Pro Gly Ala Glu Arg Gly Ala  
385 390 395 400  
Cys Pro Thr Thr Ser Gly Val Pro Ala Glu Val Glu Ala Glu Ala Pro  
405 410 415  
Asn Ser Asn Val Val Phe Ser Asn Ile Arg Phe Gly Pro Ile Gly Ser  
420 425 430  
Thr Val Ala Gly Leu Pro Gly Ala Gly Asn Gly Gly Asn Asn Gly Gly  
435 440 445  
Asn Pro Pro Pro Pro Thr Thr Ser Ser Ala Pro Ala Thr Thr  
450 455 460  
Thr Thr Ala Ser Ala Gly Pro Lys Ala Gly Arg Trp Gln Gln Cys Gly  
465 470 475 480  
Gly Ile Gly Phe Thr Gly Pro Thr Gln Cys Glu Glu Pro Tyr Thr Cys

	485	490	495
Thr Lys Leu Asn Asp Trp Tyr Ser Gln Cys Leu			
	500	505	
<210> 14			
<211> 20			
<212> DNA			
<213> Artificial Sequence			
<220>			
<223> primer			
<400> 14			
atgcgttacccg ccaagttcgc			20
<210> 15			
<211> 22			
<212> DNA			
<213> Artificial Sequence			
<220>			
<223> primer			
<400> 15			
ttacaggcac tgagagtacc ag			22